RAW SEQUENCE LISTING PATENT APPLICATION US/08/471,622A

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This Raw Listing contains the General Information Section and up to the first 5 pages.

18X1 2/04

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ENTERED
                                      SEQUENCE LISTING
1
2
           General Information:
 3
    (1)
         (i) APPLICANT: HUSE, WILLIAM D.
        (ii) TITLE OF INVENTION: SURFACE EXPRESSION LIBRARIES OF
 7
                HETEROMERIC RECEPTORS
 8
9
       (iii) NUMBER OF SEQUENCES: 76
10
11
        (iv) CORRESPONDENCE ADDRESS:
12
13
               (A) ADDRESSEE: CAMPBELL & FLORES LLP
               (B) STREET: 4370 LA JOLLA VILLAGE DRIVE, SUITE 700
14
15
               (C) CITY: SAN DIEGO
               (D) STATE: CALIFORNIA
16
               (E) COUNTRY: UNITED STATES
17
18
               (F) ZIP: 92122
19
20
         (v) COMPUTER READABLE FORM:
               (A) MEDIUM TYPE: Floppy disk
21
               (B) COMPUTER: IBM PC compatible
22
23
               (C) OPERATING SYSTEM: PC-DOS/MS-DOS
24
               (D) SOFTWARE: PatentIn Release #1.0, Version #1.25
25
26
        (vi) CURRENT APPLICATION DATA:
27
               (A) APPLICATION NUMBER: 08/471,622
28
               (B) FILING DATE: June 5, 1995
29
               (C) CLASSIFICATION:
30
      (viii) ATTORNEY/AGENT INFORMATION:
31
               (A) NAME: CAMPBELL, CATHRYN A.
32
33
               (B) REGISTRATION NUMBER: 31,815
               (C) REFERENCE/DOCKET NUMBER: P-IX 1613
34
35
        (ix) TELECOMMUNICATION INFORMATION:
36
               (A) TELEPHONE: 619-535-9001
37
               (B) TELEFAX: 619-535-8949
38
39
40
    (2) INFORMATION FOR SEQ ID NO:1:
41
42
43
         (i) SEQUENCE CHARACTERISTICS:
44
               (A) LENGTH: 7445 base pairs
               (B) TYPE: nucleic acid
45
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(C) STRANDEDNESS: both

RAW SEQUENCE LISTING PATENT APPLICATION US/08/471,622A

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(D) TOPOLOGY: circular

(xi) SEQUENCE DESCRIPTION: SEQ ID NO:1:

AATGCTACTA CTATTAGTAG AATTGATGCC ACCTTTTCAG CTCGCGCCCC AAATGAAAAT ATAGCTAAAC AGGTTATTGA CCATTTGCGA AATGTATCTA ATGGTCAAAC TAAATCTACT CGTTCGCAGA ATTGGGAATC AACTGTTACA TGGAATGAAA CTTCCAGACA CCGTACTTTA GTTGCATATT TAAAACATGT TGAGCTACAG CACCAGATTC AGCAATTAAG CTCTAAGCCA TCTGCAAAAA TGACCTCTTA TCAAAAGGAG CAATTAAAGG TACTCTCTAA TCCTGACCTG TTGGAGTTTG CTTCCGGTCT GGTTCGCTTT GAAGCTCGAA TTAAAACGCG ATATTTGAAG TCTTTCGGGC TTCCTCTTAA TCTTTTTGAT GCAATCCGCT TTGCTTCTGA CTATAATAGT CAGGGTAAAG ACCTGATTTT TGATTTATGG TCATTCTCGT TTTCTGAACT GTTTAAAGCA TTTGAGGGGG ATTCAATGAA TATTTATGAC GATTCCGCAG TATTGGACGC TATCCAGTCT AAACATTTTA CTATTACCCC CTCTGGCAAA ACTTCTTTTG CAAAAGCCTC TCGCTATTTT GGTTTTTATC GTCGTCTGGT AAACGAGGGT TATGATAGTG TTGCTCTTAC TATGCCTCGT AATTCCTTTT GGCGTTATGT ATCTGCATTA GTTGAATGTG GTATTCCTAA ATCTCAACTG ATGAATCTTT CTACCTGTAA TAATGTTGTT CCGTTAGTTC GTTTTATTAA CGTAGATTTT TCTTCCCAAC GTCCTGACTG GTATAATGAG CCAGTTCTTA AAATCGCATA AGGTAATTCA CAATGATTAA AGTTGAAATT AAACCATCTC AAGCCCAATT TACTACTCGT TCTGGTGTTT CTCGTCAGGG CAAGCCTTAT TCACTGAATG AGCAGCTTTG TTACGTTGAT TTGGGTAATG AATATCCGGT TCTTGTCAAG ATTACTCTTG ATGAAGGTCA GCCAGCCTAT GCGCCTGGTC TGTACACCGT TCATCTGTCC TCTTTCAAAG TTGGTCAGTT CGGTTCCCTT ATGATTGACC GTCTGCGCCT CGTTCCGGCT AAGTAACATG GAGCAGGTCG CGGATTTCGA CACAATTTAT CAGGCGATGA TACAAATCTC CGTTGTACTT TGTTTCGCGC TTGGTATAAT CGCTGGGGGT CAAAGATGAG TGTTTTAGTG TATTCTTTCG CCTCTTTCGT TTTAGGTTGG TGCCTTCGTA GTGGCATTAC GTATTTTACC CGTTTAATGG AAACTTCCTC ATGAAAAAGT CTTTAGTCCT CAAAGCCTCT GTAGCCGTTG CTACCCTCGT TCCGATGCTG TCTTTCGCTG CTGAGGGTGA 1.380 CGATCCCGCA AAAGCGGCCT TTAACTCCCT GCAAGCCTCA GCGACCGAAT ATATCGGTTA

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100					•		
101	TGCGTGGGCG	ATGGTTGTTG	TCATTGTCGG	CGCAACTATC	GGTATCAAGC	TGTTTAAGAA	1500
102				000.0.01.1.10	001111011100	IUIIIAOAA	1500
103	ATTCACCTCG	AAAGCAAGCT	GATAAACCGA	TACAATTAAA	GGCTCCTTTT	GGAGCCTTTT	1560
104							
105	TTTTTGGAGA	TTTTCAACGT	GAAAAAATTA	TTATTCGCAA	TTCCTTTAGT	TGTTCCTTTC	1620
106							
107	TATTCTCACT	CCGCTGAAAC	TGTTGAAAGT	TGTTTAGCAA	AACCCCATAC	AGAAAATTCA	1680
108							
109	TTTACTAACG	TCTGGAAAGA	CGACAAAACT	TTAGATCGTT	ACGCTAACTA	TGAGGGTTGT	1740
110	amamaa s sma	CM1 C1 CCCCC	mams ammam	1.0maama1.aa		EE1 000E101	
111 112	CIGIGGAAIG	CTACAGGCGT	TGTAGTTTGT	ACTGGTGACG	AAACTCAGTG	TTACGGTACA	1800
113	Τ СССТТССТΔ	TTGGGCTTGC	ТАТСССТСАА	А АТСА СССТС	<i>атаастота</i> х	CCCTCCCCCT	1860
114	ICCCIA	1100001100	INICCUIONA	AN LONGO LO	GIGGCICIGA	444144444	1000
115	TCTGAGGGTG	GCGGTTCTGA	GGGTGGCGGT	ACTAAACCTC	CTGAGTACGG	TGATACACCT	1920
116							
117	ATTCCGGGCT	ATACTTATAT	CAACCCTCTC	GACGGCACTT	ATCCGCCTGG	TACTGAGCAA	1980
118							
119	AACCCCGCTA	ATCCTAATCC	TTCTCTTGAG	GAGTCTCAGC	CTCTTAATAC	TTTCATGTTT	2040
120							
121	CAGAATAATA	GGTTCCGAAA	TAGGCAGGGG	GCATTAACTG	TTTATACGGG	CACTGTTACT	2100
122 123	CA A CCCA CTC	ACCCCGTTAA	3.3.CM@3.C@3.C	aramrarama	GMGM3 MG3 MG	1111000100	0160
124	CAAGGCACTG	ACCCCGTTAA	AACTTATTAC	CAGTACACTC	CTGTATCATC	AAAAGCCATG	2160
125	ТАТСАСССТТ	АСТЕСА АСЕС	таааттсаса	GACTGCGCTT	ТССАТТСТСС	CTTTAATGAA	2220
126			imuii i oilon	ONO TOUGOTT	1002110100	CITTATIONA	2220
127	GATCCATTCG	TTTGTGAATA	TCAAGGCCAA	TCGTCTGACC	TGCCTCAACC	TCCTGTCAAT	2280
128							
129	GCTGGCGGCG	GCTCTGGTGG	TGGTTCTGGT	GGCGGCTCTG	AGGGTGGTGG	CTCTGAGGGT	2340
130							
131	GGCGGTTCTG	AGGGTGGCGG	CTCTGAGGGA	GGCGGTTCCG	GTGGTGGCTC	TGGTTCCGGT	2400
132	a s mmmma s mm	3.MG3.3.3.G3.M	GGG3333GGGT	117112222	ami mai aaai)	
133 134	GATTTTGATT	ATGAAAAGAT	GGCAAACGCT	AATAAGGGGG	CTATGACCGA	AAATGCCGAT	2460
135	GAAAACGCGC	TACAGTCTGA	CGCTAAAGGC	ልልልሮሞሞሮልሞሞ	СТСТСССТАС	тсаттассст	2520
136			COCIMILOGO	AMOTIONII	CIGICGCIAC	IONTIACGGI	2320
137	GCTGCTATCG	ATGGTTTCAT	TGGTGACGTT	TCCGGCCTTG	CTAATGGTAA	TGGTGCTACT	2580
138							
139	GGTGATTTTG	CTGGCTCTAA	TTCCCAAATG	GCTCAAGTCG	GTGACGGTGA	TAATTCACCT	2640
140							
141	TTAATGAATA	ATTTCCGTCA	ATATTTACCT	TCCCTCCCTC	AATCGGTTGA	ATGTCGCCCT	2700
142							
143	TTTGTCTTTA	GCGCTGGTAA	ACCATATGAA	TTTTCTATTG	ATTGTGACAA	AATAAACTTA	2760
144 145	ттесетесте	TCTTTGCGTT	ጥረ ጥጥጥጥ አጥ አጥ	CTTCCC3 CCT	마까지 하스마 지하스마	A THE THE COLOR	2820
146	1100010010	1011100011	ICITITATAT	GIIGCCACCI	TIMIGIMIGI	ATTICIACG	2020
147	TTTGCTAACA	TACTGCGTAA	TAAGGAGTCT	TAATCATGCC	AGTTCTTTG	GGTATTCCGT	2880
148							
149	TATTATTGCG	TTTCCTCGGT	TTCCTTCTGG	TAACTTTGTT	CGGCTATCTG	CTTACTTTTC	2940
150							
151	TTAAAAAGGG	CTTCGGTAAG	ATAGCTATTG	CTATTTCATT	GTTTCTTGCT	CTTATTATTG	3000
152							

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						INPUL 3E1: 322//	o.raw
153 154	GGCTTAACTC	AATTCTTGTG	GGTTATCTCT	CTGATATTAG	CGCTCAATTA	CCCTCTGACT	3060
155 156	TTGTTCAGGG	TGTTCAGTTA	ATTCTCCCGT	CTAATGCGCT	TCCCTGTTTT	TATGTTATTC	3120
157 158	TCTCTGTAAA	GGCTGCTATT	TTCATTTTTG	ACGTTAAACA	AAAAATCGTT	TCTTATTTGG	3180
159 160	ATTGGGATAA	ATAATATGGC	TGTTTATTTT	GTAACTGGCA	AATTAGGCTC	TGGAAAGACG	3240
161 162	CTCGTTAGCG	TTGGTAAGAT	TCAGGATAAA	ATTGTAGCTG	GGTGCAAAAT	AGCAACTAAT	3300
163 164	CTTGATTTAA	GGCTTCAAAA	CCTCCCGCAA	GTCGGGAGGT	TCGCTAAAAC	GCCTCGCGTT	3360
165 166	CTTAGAATAC	CGGATAAGCC	TTCTATATCT	GATTTGCTTG	CTATTGGGCG	CGGTAATGAT	3420
167 168	TCCTACGATG	AAAATAAAA	CGGCTTGCTT	GTTCTCGATG	AGTGCGGTAC	TTGGTTTAAT	3480
169 170	ACCCGTTCTT	GGAATGATAA	GGAAAGACAG	CCGATTATTG	ATTGGTTTCT	ACATGCTCGT	3540
171 172	AAATTAGGAT	GGGATATTAT	TTTTCTTGTT	CAGGACTTAT	CTATTGTTGA	TAAACAGGCG	3600
173 174	CGTTCTGCAT	TAGCTGAACA	TGTTGTTTAT	TGTCGTCGTC	TGGACAGAAT	TACTTTACCT	3660
175 176	TTTGTCGGTA	CTTTATATTC	TCTTATTACT	GGCTCGAAAA	TGCCTCTGCC	TAAATTACAT	3720
177 178	GTTGGCGTTG	TTAAATATGG	CGATTCTCAA	TTAAGCCCTA	CTGTTGAGCG	TTGGCTTTAT	3780
179 180	ACTGGTAAGA	ATTTGTATAA	CGCATATGAT	ACTAAACAGG	CTTTTTCTAG	TAATTATGAT	3840
181 182	TCCGGTGTTT	ATTCTTATTT	AACGCCTTAT	TTATCACACG	GTCGGTATTT	CAAACCATTA	3900
183 184	AATTTAGGTC	AGAAGATGAA	GCTTACTAAA	ATATATTTGA	AAAAGTTTTC	ACGCGTTCTT	3960
185 186	TGTCTTGCGA	TTGGATTTGC	ATCAGCATTT	ACATATAGTT	ATATAACCCA	ACCTAAGCCG	4020
187 188		AGGTAGTCTC					4080
189 190		ATCTAAGCTA					4140
191 192		TACAGAAGCA					4200
193 194		GTAATTCAAA					4260
195 196		TCTTCTTTTG					4320
197 198						ATGTAAAAGG	4380
199 200						TCTTTATTTC	4440
201 202		GCTAATAATT					4500
203 204		AATCAGGATT					4560
205	TGATAATTCC	GCTCCTTCTG	GTGGTTTCTT	TGTTCCGCAA	AATGATAATG	TTACTCAAAC	4620

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206							
207	TTTAAAATT	AATAACGTTC	GGGCAAAGGA	TTTAATACGA	GTTGTCGAAT	TGTTTGTAAA	4680
208							
209	GTCTAATACT	TCTAAATCCT	CAAATGTATT	ATCTATTGAC	GGCTCTAATC	TATTAGTTGT	4740
210							
211	TAGTGCACCT	AAAGATATTT	TAGATAACCT	TCCTCAATTC	CTTTCTACTG	TTGATTTGCC	4800
212							
213	AACTGACCAG	ATATTGATTG	AGGGTTTGAT	ATTTGAGGTT	CAGCAAGGTG	ATGCTTTAGA	4860
214							
215	TTTTTCATTT	GCTGCTGGCT	CTCAGCGTGG	CACTGTTGCA	GGCGGTGTTA	ATACTGACCG	4920
216							
217	CCTCACCTCT	GTTTTATCTT	CTGCTGGTGG	TTCGTTCGGT	ATTTTTAATG	GCGATGTTTT	4980
218							
219	AGGGCTATCA	GTTCGCGCAT	TAAAGACTAA	TAGCCATTCA	AAAATATTGT	CTGTGCCACG	5040
220							
221	TATTCTTACG	CTTTCAGGTC	AGAAGGGTTC	TATCTCTGTT	GGCCAGAATG	TCCCTTTTAT	5100
222							
223	TACTGGTCGT	GTGACTGGTG	AATCTGCCAA	TGTAAATAAT	CCATTTCAGA	CGATTGAGCG	5160
224	mamam.						
225	TCAAAATGTA	GGTATTTCCA	TGAGCGTTTT	TCCTGTTGCA	ATGGCTGGCG	GTAATATTGT	5220
226	mamaa a ma mm	1001001100	0001 m1 0mmm	a s ammammam	100010001	ama s mamms m	F000
227 228	TCTGGATATT	ACCAGCAAGG	CCGATAGTTT	GAGTTCTTCT	ACTCAGGCAA	GTGATGTTAT	5280
229	מ א שווא א ווויט א	2022002000	OMA CA A COOM	ma ammmaaam	CAMCCACACA	CTCTTTTACT	5340
230	INCIANICAN	AGAAGIAIIG	CIACAACGGI	TAATITGCGT	GAIGGACAGA	CICITITACI	3340
231	СССТСССТС	አርጥር አጥጥአጥአ	**** * * C * C ***********************	ምር እ እር እጥጥርጥ	GGCGTACCGT	TCCTGTCTAA	5400
232	0001000010	ACTORITATA	MANACACTIC	ICANOATICI	GGCGTACCGT	ICCIGICIAA	3400
233	AATCCCTTTA	ATCGGCCTCC	TGTTTAGCTC	CCGCTCTGAT	TCCAACGAGG	AAAGCACGTT	5460
234							
235	ATACGTGCTC	GTCAAAGCAA	CCATAGTACG	CGCCCTGTAG	CGGCGCATTA	AGCGCGGCGG	5520
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237	GTGTGGTGGT	TACGCGCAGC	GTGACCGCTA	CACTTGCCAG	CGCCCTAGCG	CCCGCTCCTT	5580
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239	TCGCTTTCTT	CCCTTCCTTT	CTCGCCACGT	TCGCCGGCTT	TCCCCGTCAA	GCTCTAAATC	5640
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244							
245	CGTTGGAGTC	CACGTTCTTT	AATAGTGGAC	TCTTGTTCCA	AACTGGAACA	ACACTCAACC	5820
246	*********						
247	CTATCTCGGG	CTATTCTTT	GATTTATAAG	GGATTTTGCC	GATTTCGGAA	CCACCATCAA	5880
248 249	3.03.003.mmmm	оодотаатаа	0003330030	CGTGGACCGC	mmaamaa sa	mamama » aaa	E040
250	ACAGGATTTT	CGCCTGCTGG	GGCAAACCAG	CGTGGACCGC	TTGCTGCAAC	TCTCTCAGGG	5940
250	селессени	A A CCCCC A A TCC	A C C T C T T T C C C	CGTCTCGCTG	CTC A A A A C A A	3 3 3 CC 3 CC CC	6000
252	CCAGGCGGIG	ANGUUMATC	MOCIGIIGCC	CGICICGCIG	GIGAAAAGAA	MMMCCMCCCI	0000
253	GGCGCCCAAT	ACGCAAACCG	ССТСТССССС	CGCGTTGGCC	ሪያ ተ መረያ ተመል አ	тасъвствае	6060
254	JJCCCCARI	oonnood	231010000	2300110000	CALLCALIAN	100001000	3000
255	ACGACAGGTT	TCCCGACTGG	AAAGCGGGCA	GTGAGCGCAA	CGCAATTAAT	GTGAGTTAGC	6120
256						 •	-
257	TCACTCATTA	GGCACCCCAG	GCTTTACACT	TTATGCTTCC	GGCTCGTATG	TTGTGTGGAA	6180
258							

SEQUENCE VERIFICATION REPORT PATENT APPLICATION US/08/471,622A

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